

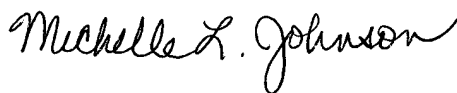
- II. Claims 7-12 and 15 drawn to a polynucleotide encoding a carboxypeptidase having amino acid sequence of SEQ ID NO:2 from at least position 19 through position 738, to vectors or recombinant viruses comprising the polynucleotide, to host cells comprising the polynucleotides, to a composition of making a first encoded carboxypeptidase in a host cell using the polynucleotide, classified, inter alia, in class 536, subclass 23.2.
- III. Claims 143 and 14 drawn to antibodies, classified in class 530, subclass 387.1.
- IV. Claim 16, drawn to an alternate method of use of a polynucleotide encoding the amino acid sequence of SEQ ID NO:2 from at least position 19 through position 738 in a method of detecting gene expression, classified in class 435, subclass 6.
- V. Claims 1-3 and 17, drawn, in part, to a carboxypeptidase having the amino acid sequence of SEQ ID NO:6 from at least position 19 through position 738, classified in class 435, subclass 226.
- VI. Claims 1-3 and 17, drawn, in part, to a carboxypeptidase having the amino acid sequence of SEQ ID NO:9 from at least position 19 through position 738, classified in class 435, subclass 226.
- VII. Claims 18-20 drawn, in part to a polynucleotide encoding a carboxypeptidase having the amino acid sequence of SEQ ID NO: 6 from at least position 19 through position 738 to vectors comprising the polynucleotide, and to host cells comprising the polynucleotide, classified inter alia in class 536, subclass 23.2.
- VIII. Claims 18-20 drawn, in part to a polynucleotide encoding a carboxypeptidase having the amino acid sequence of SEQ ID NO: 6 from at least position 19 through position 738 to vectors comprising the polynucleotide, and to host cells comprising the polynucleotide, classified inter alia in class 536, subclass 23.2.

In response to the Requirement for Restriction Election, Applicant elects with traverse Groups II, VII, and VIII, claims 7-12, 15, and 18-20. These three groups are drawn to polynucleotide sequences, as well as vectors, recombinant viruses, host cells comprising the polynucleotide sequences, and compositions comprising the vectors, as well as a method of making the carboxypeptidase using the polynucleotide sequences. These groups are properly examined together because the three sequences (the nucleotide equivalents of SEQ ID NOS 2, 6, and 9 or fragments thereof) have significant structural similarity. This similarity means that a search for one sequence will result in the identification of sequences related to all three. Accordingly, the examination of the three restriction groups together is proper. In a telephone conversation on February 27, 2003, Examiner Moore confirmed this approach. Applicant respectfully thanks the Examiner for providing his views on the election requirement.

Claims 1-6, 14, 16, and 17 are withdrawn from the case; however, Applicant reserves the right to pursue and prosecute these claims in continuation and divisional applications.

Early consideration and allowance of the pending claims is respectfully requested. If the Patent Examiner believes that a telephone interview would expedite prosecution of the patent application, please call the undersigned at (206) 442-6627.

Respectfully Submitted,



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Enclosures:

Petition and Fee for Extension of Time (in duplicate)
Amendment Fee Transmittal (in duplicate)
Associate Power of Attorney
Postcard